

**Basel Biometric Society**  
**Workshop/Seminar, 2024**

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*FIRST ANNOUNCEMENT*

**Controlling the chances of  
false discoveries in exploratory  
analysis of clinical trials**

**Date: 29th August 2024**

**Time: 14:00-17:00 CET**

**Venue: Virtual**

**Costs: Free**

**Registration: [LINK](#)**

Note: Dial-in details will be communicated to registered participants

**Organizers:** Kostas Sechidis (Novartis), Frank Bretz (Novartis)

While the primary focus of clinical trials is to estimate causal effects, the collected data can also be invaluable for additional research, such as identifying variables and/or groups of patients with desirable characteristics. Some common exploratory analysis activities focus on using clinical trial data for variable selection. For example, we may want to identify baseline variables that are strongly associated with the disease outcome, irrespective of the treatment assignment (i.e., prognostic variables) or baseline variables that influence the treatment effect (i.e., predictive variables). Clinical trial data can also be used for subgroup discovery, where, for example, we aim to identify groups of patients that experience a significant treatment effect. In all these selection problems, it is critical to control the chances of false discoveries (type-I errors) to provide guarantees concerning the replicability of our results. The focus of this session is on recent methodologies for performing this type of selection by controlling the type-I error rate.

We hope that many of you can attend this exciting seminar!

→ See the next page for the Program

## Program

14.00-14.15	<b>Welcome remarks and setting the scene</b>	Dominic Magirr (Novartis, BBS board member) and Frank Bretz (Novartis)
<b>Part 1: Controlled variable selection</b>		
14.15-14.40	<b>Using knockoffs for type-I error controlled prognostic and predictive variable selection</b>	Kostas Sechidis (Novartis)
14.40-15.05	<b>Statistical Interpretation of High-Dimensional Prediction Models using Conditional Permutation Importance</b>	Denis Engemann (Roche)
15.05-15.30	<b>Controlled Discovery and Localization of Signals via Bayesian Linear Programming</b>	Asher Spector (Stanford University)
15.30-15.45	<b>Break</b>	
<b>Part 2: Controlled subgroup discovery</b>		
15.45-16.10	<b>Overview of modern approaches for identifying and evaluating heterogeneous treatment effects from clinical trials</b>	Ilya Lipkovich (Elli Lilly)
16.10-16.35	<b>Subgroup selection with strong type-I error control under monotonicity constraints</b>	Manuel M. Müller (University of Cambridge)
16.35-17.00	<b>Interactive identification of individuals with positive treatment effect while controlling false discoveries</b>	Aaditya Ramdas (Carnegie Mellon University)